## SEQUENCE LISTING

<110>	Brugliera, Filippa Fanaka, Yoshikazu Mason, John	
<120>	Flavonoid 3',5' Hydroxylase Gene Sequences and Uses Therefor	
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Gly Ala Ile Met Tyr Leu Lys Val Gly Thr Cys Gly Met Ala Val Ala 65 70 75 80

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Gly Ala Ile Met Tyr Leu Lys Val Gly Thr Cys Gly Met Val Val Ala 65 70 75 80

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Cys Glu Ala Ser Arg Cys Gly Glu Pro Val Val Leu Ala Glu Met Leu 165 170 175

Thr Tyr Ala Met Ala Asn Met Ile Gly Gln Val Ile Leu Ser Arg Arg 180 185 190

Val Phe Val Thr Lys Gly Thr Glu Ser Asn Glu Phe Lys Asp Met Val 195 200 205

Val Glu Leu Met Thr Ser Ala Gly Tyr Phe Asn Ile Gly Asp Phe Ile 210 215 220

Pro Ser Ile Ala Trp Met Asp Leu Gln Gly Ile Glu Arg Gly Met Lys 225 230 235 240

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tgaaaataat gaatcggaaa aaaaaaaaaa aaaaaaaaaa
<210> 12 <211> 506 <212> PRT <213> viola
<400> 12
Met Ala Ile Leu Val Thr Asp Phe Val Val Ala Ala Ile Ile Phe Leu 1 5 10 15
Ile Thr Arg Phe Leu Val Arg Ser Leu Phe Lys Lys Pro Thr Arg Pro 20 25 30
Leu Pro Pro Gly Pro Leu Gly Trp Pro Leu Val Gly Ala Leu Pro Leu 35 40 45
Leu Gly Ala Met Pro His Val Ala Leu Ala Lys Leu Ala Lys Lys Tyr 50 55 60
Gly Pro Ile Met His Leu Lys Met Gly Thr Cys Asp Met Val Val Ala 65 70 75 80
Ser Thr Pro Glu Ser Ala Arg Ala Phe Leu Lys Thr Leu Asp Leu Asn 85 90 95
Phe Ser Asn Arg Pro Pro Asn Ala Gly Ala Ser His Leu Ala Tyr Gly 100 105 110
Ala Gln Asp Leu Val Phe Ala Lys Tyr Gly Pro Arg Trp Lys Thr Leu 115 120 125
Arg Lys Leu Ser Asn Leu His Met Leu Gly Gly Lys Ala Leu Asp Asp 130 135 140
Trp Ala Asn Val Arg Val Thr Glu Leu Gly His Met Leu Lys Ala Met 145 150 155 160
Cys Glu Ala Ser Arg Cys Gly Glu Pro Val Val Leu Ala Glu Met Leu 165 170 175

Thr Tyr Ala Met Ala Asn Met Ile Gly Gln Val Ile Leu Ser Arg Arg 180 185 190

Val Phe Val Thr Lys Gly Thr Glu Ser Asn Glu Phe Lys Asp Met Val 195 200 205

Val Glu Leu Met Thr Ser Ala Gly Tyr Phe Asn Ile Gly Asp Phe Ile 210 215 220

Pro Ser Ile Ala Trp Met Asp Leu Gln Gly Ile Glu Arg Gly Met Lys 225 230 235 240

Lys Leu His Thr Lys Phe Asp Val Leu Leu Thr Lys Met Val Lys Glu 245 250 255

His Arg Ala Thr Ser His Glu Arg Lys Gly Lys Ala Asp Phe Leu Asp 260 265 270

Val Leu Leu Glu Glu Cys Asp Asn Thr Asn Gly Glu Lys Leu Ser Ile 275 280 285

Thr Asn Ile Lys Ala Val Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp 290 295 300

Thr Ser Ser Ser Ile Ile Glu Trp Ala Leu Thr Glu Met Ile Lys Asn 305 310 315 320

Pro Thr Ile Leu Lys Lys Ala Gln Glu Glu Met Asp Arg Val Ile Gly 325 330 335

Arg Asp Arg Leu Leu Glu Ser Asp Ile Ser Ser Leu Pro Tyr Leu 340 345 350

Gln Ala Ile Ala Lys Glu Thr Tyr Arg Lys His Pro Ser Thr Pro Leu 355 360 365

Asn Leu Pro Arg Ile Ala Ile Gln Ala Cys Glu Val Asp Gly Tyr Tyr 370 375 380

Ile Pro Lys Asp Ala Arg Leu Ser Val Asn Ile Trp Ala Ile Gly Arg 385 390 395 400

Asp Pro Asn Val Trp Glu Asn Pro Leu Glu Phe Leu Pro Glu Arg Phe
405 410 415

Leu Ser Glu Glu Asn Gly Lys Ile Asn Pro Gly Gly Asn Asp Phe Lys

420 425 430

Leu Ile Pro Phe Gly Ala Gly Arg Arg Ile Cys Ala Gly Thr Arg Met 435 440 445

Gly Met Val Leu Val Ser Tyr Ile Leu Gly Thr Leu Val His Ser Phe 450 455 460

Asp Trp Lys Leu Pro Asn Gly Val Ala Glu Leu Asn Met Asp Glu Ser 465 470 475 480

Phe Gly Leu Ala Leu Gln Lys Ala Val Pro Leu Ser Ala Leu Val Ser 485 490 495

Pro Arg Leu Ala Ser Asn Pro Tyr Ala Thr 500 505

<210> 13 <211> 1659 <212> DNA

<213> salvia

<400> 13

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tattaaagcc cttttactga atttatttac tgcagqqaca qacacatcqt cqaqcataat 960 agaatgggca ttagcggaga tgatcaagaa tccaagcatc caaaaaaqqq ctcaccaaqa 1020 gatggacaga gtcatcggga gagagcggcg tttgctcgaa tccgacatcc caaatctqcc 1080 atacctcaaa gccatatgca aagaggcata ccgaaaacac ccttccacgc cactaaacct 1140 gcctcggatc tccacggatg catgcgtcgt cgatggctac cacatcccca agaacacgag 1200 gttgagcgtc aacatctggg ccataggccg agatcccgac gtttgggaga atccccttga 1260 cttcaaccct gacaggttta tgtcagggtt gcaggggatt gagcccggag ggaatcactt 1320 cgagctcatt ccctttgggg cggggcgcag gatctgcgcc ggcagcagaa tggggattgt 1380 aatagtggag tatttgctgg cgacactcgt gcactctttc gaatgggatt tgccggccgg 1440 ctcagcggag atggacatgg aggaggtgtt cgggctqqcc ttqcaqaaaq ctqtaccact 1500 tgctgctagg ctcactccta ggttgccttc acattgctat gcacctcctt ctatttaatt 1560 tgcatattta catatgttgt gttacattga gcctttqcat atqttqtatc caacctatct 1620 tataacttgt gcatgaaatt gaaaaaaaaa aaaaaaaaa 1659

<210> 14

<211> 520

<212> PRT

<213> salvia

<400> 14

Gly Thr Ser Met Glu Ala Gln Glu Asn Met Leu Leu Ile Ala Arg Ala 1 5 10 15

Leu Val Val Ala Ser Leu Leu Tyr Ile Leu Ile Arg Met Phe Ile Ser 20 25 30

Lys Leu Ser Thr Thr Gly His Pro Leu Pro Pro Gly Pro Arg Gly Phe 35 40 45

Leu Val Val Gly Ser Leu Pro Leu Leu Gly Asp Met Pro His Val Ala 50 55 60

Leu Ala Lys Met Ala Lys Thr Tyr Gly Pro Ile Met Tyr Leu Lys Met 65 70 75 80

Gly Thr Val Gly Met Val Val Ala Ser Thr Pro Asp Ala Ala Arg Ala 85 90 95

Phe Leu Lys Thr His Asp Ala Asn Phe Ser Asn Arg Pro Val Asn Ala

100 105 110

Gly	Ala	Thr	Ile	Leu	Ala	Tyr	Asn	Ala	Gln	Asp	Met	Val	Phe	Ala	Pro
		115					120					125			

- Tyr Gly Pro Lys Trp Arg Leu Leu Arg Lys Leu Ser Ser Leu His Met 130 135 140
- Leu Gly Ser Lys Ala Leu Glu Glu Trp Ala Asp Val Arg Thr Ser Glu 145 150 155 160
- Val Gly His Met Leu Ala Ala Met His Glu Ala Ser Arg Leu Gly Glu 165 170 175
- Ala Val Gly Leu Pro Glu Met Leu Val Tyr Ala Thr Ala Asn Met Ile 180 185 190
- Gly Gln Val Ile Leu Ser Arg Arg Val Phe Val Thr Lys Gly Lys Glu 195 200 205
- Met Asn Glu Phe Lys Glu Met Val Val Glu Leu Met Thr Thr Ala Gly 210 215 220
- Tyr Phe Asn Ile Gly Asp Phe Ile Pro Trp Leu Ala Trp Met Asp Leu 225 230 235 240
- Gln Gly Ile Glu Arg Gly Met Lys Lys Leu His Lys Lys Trp Asp Arg 245 250 255
- Leu Ile Gly Lys Met Leu Asp Asp Arg Leu Lys Ser Thr Tyr Lys Arg 260 265 270
- Asn Asp Lys Pro Asp Leu Leu Asp Ser Leu Leu Ala Asn His Asp Asp 275 280 285
- Glu Ser Lys Asp Asp Asp Glu Asp Cys Lys Leu Thr Thr Thr Asn Ile 290 295 300
- Lys Ala Leu Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser 305 310 315 320
- Ser Ile Ile Glu Trp Ala Leu Ala Glu Met Ile Lys Asn Pro Ser Ile 325 330 335

Gln	Lys	Arg	Ala 340	His	Gln	Glu	Met	Asp 345	Arg	Val	Ile	Gly	Arg 350	Glu	Arg					
Arg	Leu	Leu 355	Glu	Ser	Asp	Ile	Pro 360	Asn	Leu	Pro	Tyr	Leu 365	Lys	Ala	Ile					
Сув	Lys 370	Glu	Ala	Tyr	Arg	Lys 375	His	Pro	Ser	Thr	Pro 380	Leu	Asn	Leu	Pro					
Arg 385	Ile	Ser	Thr	Asp	Ala 390	Cys	Val	Val	Asp	Gly 395	Tyr	His	Ile	Pro	Lys 400					
Asn	Thr	Arg	Leu	Ser 405	Val	Asn	Ile	Trp	Ala 410	Ile	Gly	Arg	Asp	Pro 415	Asp					
Val	Trp	Glu	Asn 420	Pro	Leu	Asp	Phe	Asn 425	Pro	Asp	Arg	Phe	Met 430	Ser	Gly					
Leu	Gln	Gly 435	Ile	Glu	Pro	Gly	Gly 440	Asn	His	Phe	Glu	Leu 445	Ile	Pro	Phe					
Gly	Ala 450	Gly	Arg	Arg	Ile	Cys 455	Ala	Gly	Ser	Arg	Met 460	Gly	Ile	Val	Ile					
Val 465	Val Glu Tyr Leu Leu Ala Thr Leu Val His Ser Phe Glu Trp Asp Leu 465 470 475 480																			
Pro	Pro Ala Gly Ser Ala Glu Met Asp Met Glu Glu Val Phe Gly Leu Ala 485 490 495																			
Leu	Gln	Lys	Ala 500	Val	Pro	Leu	Ala	Ala 505	Arg	Leu	Thr	Pro	Arg 510	Leu	Pro					
Ser	His	Cys 515	Tyr	Ala	Pro	Pro	Ser 520													
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cat	cctta	act o	ctaca	atttt	g at	ccgt	atgt	tta	atcto	caaa	cateettact ctacattttg atcegtatgt ttateteaaa attgageace eeeggeeace									

ctctgccccc ggggccgagg gqctttccaq tqqtqqqctc ccttcccttq ctqqqcqaca 180 tgccacatgt tgccctagca aaaatggcca aaacttatgg cccqatcatg tacttqaaaa 240 tgggcacagt cggcatggtc gtggcgtcca cgccagacgc ggcgcgggcg ttcctaaaaa 300 cccaggacgc taatttctct aaccggccgg tcaacgcggg tgccaccatc ctggcataca 360 atgcccagga catggtgttt gccccgtacg gccccaagtg gagattgctg aggaagctga 420 gcagtctcca catgctgggg agcaaggccc tggaggagtg ggccgacgtc cggacctcgg 480 aggtggggca catgctggcg gcgatgcacg aggccagccg cctggacgag gccgtggggt 540 tgccggagat gctggtgtac gcgacggcga acatgatcgg gaaggtgata ttgagccgga 600 gagttttcgt gacgaaaggg aaggagatga atgagttcaa ggaaatggtq qtqqaqctca 660 tgaccacage tggctattte aacattggtg attteattee atggettget tggatggatt 720 tgcaggggat tgagagaggg atgaagaaac tgcacaagaa gtgggaccgc ttgatcggta 780 agatgctgga tgatcgattg aaatcaacct acaaacgcaa cgacaagcca gatcttcttg 840 attetetett ggeaaateat gatgatgaga gtaaggatga tgatgaggat tgeaagetea 900 ccaccaccaa tattaaagcc cttttactga atttatttac tgcagggaca gacacatcgt 960 cgagcataat agaatgggca ctagcggaga tgatcaagaa tccaagcatc caaaaaaggg 1020 ctcaccaaga gatggacaga gtcatcggga gagagcggcg tttgctcgaa tccgacatcc 1080 caaatctgcc atacctcaaa gccatatgca aagaggcata ccgaaaacac ccttccacgc 1140 cactaaacct gcctcggatc tccacggatg catgcgtcgt cgatggctac cacatcccca 1200 agaacacgag gttgagcgtc aacatctggg ccataggccg agatcccgac qtttqqqaqa 1260 atccccttga cttcaaccct gacaggttta tqtcaqqqtt qcaqqqqatt qaqcccqqaq 1320 ggaatcactt cgagctcatt ccctttgggg cggggcgcag gatctgcgcc ggcagcagaa 1380 tggggattgt aatagtggag tatttgctgg cgacactcgt gcactctttc gaatgggatt 1440 tgccagccgg ctcagcggag atggacatgg aggaggtgtt cgggctggcc ttgcagaaaq 1500 ctgtaccact tgctgctagg ctcactccta ggttgccttc acattgctat gcacctcctt 1560 ctatttaatt tgcatattta tatatgttgt gttacattga aaaaaaaaa aaaaaaa 1617

Met Glu Ala Gln Glu Asn Met Leu Leu Ile Ala Arg Ala Leu Val Val

<sup>&</sup>lt;210> 16 <211> 518 <212> PRT <213> salvia

<sup>&</sup>lt;400> 16

Ala	Ser	Leu	Leu	Tyr	Ile	Leu	Ile	Arg	Met	Phe	Ile	Ser	Lys	Leu	Ser
			20					25					30		

1

Thr Pro Gly His Pro Leu Pro Pro Gly Pro Arg Gly Phe Pro Val Val
35 40 45

Gly Ser Leu Pro Leu Leu Gly Asp Met Pro His Val Ala Leu Ala Lys 50 60

Met Ala Lys Thr Tyr Gly Pro Ile Met Tyr Leu Lys Met Gly Thr Val 65 70 75 80

Gly Met Val Val Ala Ser Thr Pro Asp Ala Ala Arg Ala Phe Leu Lys 85 90 95

Thr Gln Asp Ala Asn Phe Ser Asn Arg Pro Val Asn Ala Gly Ala Thr 100 105 110

Ile Leu Ala Tyr Asn Ala Gln Asp Met Val Phe Ala Pro Tyr Gly Pro 115 120 125

Lys Trp Arg Leu Leu Arg Lys Leu Ser Ser Leu His Met Leu Gly Ser 130 135 140

Lys Ala Leu Glu Glu Trp Ala Asp Val Arg Thr Ser Glu Val Gly His 145 150 155 160

Met Leu Ala Ala Met His Glu Ala Ser Arg Leu Asp Glu Ala Val Gly 165 170 175

Leu Pro Glu Met Leu Val Tyr Ala Thr Ala Asn Met Ile Gly Lys Val 180 185 190

Ile Leu Ser Arg Arg Val Phe Val Thr Lys Gly Lys Glu Met Asn Glu
195 200 205

Phe Lys Glu Met Val Val Glu Leu Met Thr Thr Ala Gly Tyr Phe Asn 210 215 220

Ile Gly Asp Phe Ile Pro Trp Leu Ala Trp Met Asp Leu Gln Gly Ile 225 230 235 240

Glu Arg Gly Met Lys Lys Leu His Lys Lys Trp Asp Arg Leu Ile Gly 245 250 Lys Met Leu Asp Asp Arg Leu Lys Ser Thr Tyr Lys Arg Asn Asp Lys 260 265 Pro Asp Leu Leu Asp Ser Leu Leu Ala Asn His Asp Asp Glu Ser Lys Asp Asp Asp Glu Asp Cys Lys Leu Thr Thr Thr Asn Ile Lys Ala Leu 290 295 Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser Ser Ile Ile Glu Trp Ala Leu Ala Glu Met Ile Lys Asn Pro Ser Ile Gln Lys Arq Ala His Gln Glu Met Asp Arg Val Ile Gly Arg Glu Arg Arg Leu Leu Glu Ser Asp Ile Pro Asn Leu Pro Tyr Leu Lys Ala Ile Cys Lys Glu Ala Tyr Arg Lys His Pro Ser Thr Pro Leu Asn Leu Pro Arg Ile Ser 370 375 Thr Asp Ala Cys Val Val Asp Gly Tyr His Ile Pro Lys Asn Thr Arg 390 Leu Ser Val Asn Ile Trp Ala Ile Gly Arg Asp Pro Asp Val Trp Glu 405 Asn Pro Leu Asp Phe Asn Pro Asp Arg Phe Met Ser Gly Leu Gln Gly 420 425 Ile Glu Pro Gly Gly Asn His Phe Glu Leu Ile Pro Phe Gly Ala Gly 435 440 Arg Arg Ile Cys Ala Gly Ser Arg Met Gly Ile Val Ile Val Glu Tyr 455 Leu Leu Ala Thr Leu Val His Ser Phe Glu Trp Asp Leu Pro Ala Gly

470

Ser Ala Glu Met Asp Met Glu Glu Val Phe Gly Leu Ala Leu Gln Lys 485 490 495

Ala Val Pro Leu Ala Ala Arg Leu Thr Pro Arg Leu Pro Ser His Cys 500 505 510

Tyr Ala Pro Pro Ser Ile 515

<210> 17

<400> 17

<211> 1730 <212> DNA

<213> sollya

<220>

<221> misc feature

<222> (1372)..(1372)

<223> n = any nucleotide

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tatagetgag ettataegea ateetaaaat ettageeeaa geeeaaagag agttggaett

ggtggttggt ccaaatagac ttgtaacgga tttqqacctc aaacaattaa cctacctaca 1080 agccatcgtc aaagaaacct ttcggctaca tcctgctacc ccactttcac ttccacggat 1140 cgcaaccgaa agctgtgaaa tcaacgggtt ttacattcca aagggctcaa cacttctcgt 1200 taacatatgg gccataggcc gtgatccaaa cacttgggct gaaccattgg tattccgacc 1260 tgaacgattc ttatcggatg gtgaaagtcc taatgttgat gttaaaggac gtaattttga 1320 attgatacca tttggggcgg ggcgaagaat ttgtgctggg atgaactttg gnctacgcat 1380 ggtccagtta gttactgcaa cgttaattca tgcatttaac tgggagttgc cagaagggga 1440 attgccagaa aatatgaata tggaggaaga ctatgggatt agcttgcaac ggacagtgcc 1500 attagttgtt catccaaagc ccagactaga ccatgaagtt tatcagtccc atggagttgt 1560 aaactqaqta cattcatqaa ctqacccaqa aqctqtcaqa tqtcqtctta tattqcctta 1620 tgtagtgcga cccttgtgtg ttttttatgt attgttttgt acaaggttga agcccqtgcq 1680 1730

<210> 18

<211> 521

<212> PRT

<213> sollya

<400> 18

Met Ala Thr Thr Leu Glu Phe Ile Leu Cys Phe Thr Ile Thr Ala Leu
1 5 10 15

Pro Phe Leu Tyr Cys Ile Leu Asn Met Arg Ile Leu Leu Asn Arg His 20 25 30

Pro Arg Ser Leu Pro Pro Gly Pro Arg Pro Trp Pro Ile Val Gly Asn 35 40 45

Leu Pro His Leu Gly Thr Lys Pro His His Ser Ile Ala Ala Met Ala 50 55 60

Arg Lys Tyr Gly Pro Leu Leu His Leu Arg Met Gly Ile Val His Val 65 70 75 80

Val Val Ala Ala Ser Ala Asp Val Ala Ala Gln Phe Leu Lys Asn Asp 85 90 95

Ala Asn Phe Ser Ser Arg Pro Pro Asn Ser Gly Ala Lys His Met Ala
100 105 110

Tyr Asn Tyr His Asp Met Val Phe Ala Pro Tyr Gly Pro Arg Trp Arg Met Leu Arg Lys Ile Cys Ala Leu His Ile Phe Ser Ala Lys Ala Leu Asp Asp Phe His Arg Val Arg Glu Glu Val Ala Ile Leu Ala Arg 150 155 Thr Leu Ala His Ala Gly Gln Lys Pro Val Asn Leu Gly Gln Leu Phe 165 170 Ser Thr Cys Asn Ala Asn Ala Leu Ser Val Leu Met Leu Gly Arg Arg 185 Leu Phe Ser Thr Glu Val Asp Ser Lys Ala Tyr Asp Phe Lys Gln Met 195 200 Val Val Glu Leu Met Thr Leu Ala Gly Glu Phe Asn Val Ser Asp Phe 210 215 220 Ile Pro Pro Leu Glu Trp Leu Asp Leu Gln Gly Val Ala Ala Lys Met 225 230 235 240 Lys Asn Val His Asn Arg Phe Asp Ala Phe Leu Asn Val Ile Leu Glu 245 250 Glu His Lys Leu Lys Leu Asn Asn Ser Gly His Gly Glu Gln Lys His 260 Met Asp Leu Leu Ser Thr Leu Ile Leu Leu Lys Asp Asp Ala Asp Ser 275 Glu Gly Gly Lys Leu Thr Asp Thr Glu Ile Lys Ala Leu Leu Asn 290 Leu Phe Ser Ala Gly Thr Asp Thr Ser Ser Ser Thr Ile Glu Trp Val 310 Ile Ala Glu Leu Ile Arg Asn Pro Lys Ile Leu Ala Gln Ala Gln Arg 325

Glu Leu Asp Leu Val Val Gly Pro Asn Arg Leu Val Thr Asp Leu Asp

340 345 350

Leu Lys Gln Leu Thr Tyr Leu Gln Ala Ile Val Lys Glu Thr Phe Arg 355 360 365

Leu His Pro Ala Thr Pro Leu Ser Leu Pro Arg Ile Ala Thr Glu Ser 370 375 380

Cys Glu Ile Asn Gly Phe Tyr Ile Pro Lys Gly Ser Thr Leu Leu Val 385 390 395 400

Asn Ile Trp Ala Ile Gly Arg Asp Pro Asn Thr Trp Ala Glu Pro Leu 405 410 415

Val Phe Arg Pro Glu Arg Phe Leu Ser Asp Gly Glu Ser Pro Asn Val 420 425 430

Asp Val Lys Gly Arg Asn Phe Glu Leu Ile Pro Phe Gly Ala Gly Arg 435 440 445

Arg Ile Cys Ala Gly Met Asn Phe Gly Leu Arg Met Val Gln Leu Val 450 455 460

Thr Ala Thr Leu Ile His Ala Phe Asn Trp Glu Leu Pro Glu Gly Glu 465 470 475 480

Leu Pro Glu Asn Met Asn Met Glu Glu Asp Tyr Gly Ile Ser Leu Gln
485 490 495

Arg Thr Val Pro Leu Val Val His Pro Lys Pro Arg Leu Asp His Glu 500 505 510

Val Tyr Gln Ser His Gly Val Val Asn 515 520

<210> 19

<211> 37

<212> DNA

<213> petunia

<400> 19

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<210> 20

<211> 1748

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aaaaaaaa

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Cys Ile Pro Arg Gln Arg Val Arg Asn Arg Thr Ser Leu Arg Thr Trp 210 215 220

Trp Phe Lys Leu Met Thr Val Ala Gly Tyr Phe Asn Ile Gly Asp Phe 225 230 235 240

Phe Pro Phe Leu Ala Arg Arg Arg Gln Gly Ile Glu Arg Gly Met 245 250 255

Lys Thr Leu His Asn Lys Lys Asp Asp Leu Leu Thr Thr Met Ile His 260 265 270

Glu His Val Ala Ser Ala His Lys Arg Lys Gly Lys Pro Pro Phe Leu 275 280 285

Asp Val Leu Met Ala His His Thr Asn Glu Ser His Glu Leu Ser Leu 290 295 300

Thr Asn Ile Lys Ala Leu Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp 305 310 315 320

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Gln Ala Ile Cys Lys Glu Thr Tyr Arg Lys His Pro Ser Thr Pro Leu 370 375 380

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Pro Ile Leu Gly Ala Leu Pro Leu Gly Asn Met Pro His Val Thr 50 55 60

Phe Ala Asn Met Ala Lys Lys Tyr Gly Ser Val Met Tyr Leu Lys Val 65 70 75 80

Gly Ser His Gly Leu Ala Ile Ala Ser Thr Pro Asp Ala Ala Lys Ala Phe Leu Lys Thr Leu Asp Leu Asn Phe Ser Asn Arg Pro Pro Asn Ala 105 Gly Ala Thr His Leu Ala Tyr Asn Ala Gln Asp Met Val Phe Ala His 120 Tyr Gly Pro Lys Trp Lys Leu Leu Arg Lys Leu Ser Asn Leu His Met 135 Leu Gly Gly Lys Ala Leu Glu Asn Trp Ala Asp Val Arg Lys Thr Glu 150 Leu Gly Tyr Met Leu Lys Ala Met Phe Glu Ser Ser Gln Asn Asn Glu 165 170 Pro Val Met Ile Ser Glu Met Leu Thr Tyr Ala Met Ala Asn Met Leu 180 185 Ser Gln Val Ile Leu Ser Arg Arg Val Phe Asn Lys Lys Gly Ala Lys 195 200 Ser Asn Glu Phe Lys Asp Met Val Val Glu Leu Met Thr Ser Ala Gly 210 215 220 Tyr Phe Asn Ile Gly Asp Phe Ile Pro Ser Ile Gly Trp Met Asp Leu 225 230 235 Gln Gly Ile Glu Gly Gly Met Lys Arg Leu His Lys Lys Phe Asp Val 245 Leu Leu Thr Arg Leu Leu Asp Asp His Lys Arg Thr Ser Gln Glu Arg 260 Lys Gln Lys Pro Asp Phe Leu Asp Phe Val Ile Ala Asn Gly Asp Asn 275 Ser Asp Gly Glu Arg Leu Asn Thr Asp Asn Ile Lys Ala Leu Leu Leu 290 Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser Ser Ile Ile Glu Trp 305

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Arg Lys His Pro Ser Thr Pro Leu Asn Leu Pro Arg Asn Cys Ile Arg 370 375

Gly His Val Asp Val Asn Gly Tyr Tyr Ile Pro Lys Gly Thr Arg Leu

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Lys Ile Asp Pro Arg Gly Asn His Phe Glu Leu Ile Pro Phe Gly Ala 435 440

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Asn Met Val Val Ala Ser Thr Pro Ala Ala Ala Arg Ala Phe Leu Lys 85 90 95

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Met Ala Ser Leu Thr Asp Ile Ala Ala Ile Arg Glu Ala Gln Arg Ala 1 5 10 15

Gln Gly Pro Ala Thr Ile Leu Ala Ile Gly Thr Ala Thr Pro Ala Asn 20 25 30

<sup>&</sup>lt;210> 29

<sup>&</sup>lt;211> 398

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> chrysanthemum

<sup>&</sup>lt;400> 29

Cys Val Tyr Gln Ala Asp Tyr Pro Asp Tyr Tyr Phe Arg Ile Thr Lys 35 40 Ser Glu His Met Val Asp Leu Lys Glu Lys Phe Lys Arg Met Cys Asp Lys Ser Met Ile Arg Lys Arg Tyr Met His Leu Thr Glu Glu Tyr Leu 70 Lys Glu Asn Pro Asn Leu Cys Glu Tyr Met Ala Pro Ser Leu Asp Ala Arg Gln Asp Val Val Val Glu Val Pro Lys Leu Gly Lys Glu Ala Ala Thr Lys Ala Ile Lys Glu Trp Gly Gln Pro Lys Ser Lys Ile Thr His Leu Ile Phe Cys Thr Thr Ser Gly Val Asp Met Pro Gly Ala Asp Tyr Gln Leu Thr Lys Leu Leu Gly Leu Arg Pro Ser Val Lys Arg Phe 150 Met Met Tyr Gln Gln Gly Cys Phe Ala Gly Gly Thr Val Leu Arg Leu 165 Ala Lys Asp Leu Ala Glu Asn Asn Lys Asp Ala Arg Val Leu Val Val 180 Cys Ser Glu Ile Thr Ala Val Thr Phe Arg Gly Pro Asn Asp Thr His Leu Asp Ser Leu Val Gly Gln Ala Leu Phe Gly Asp Gly Ala Ala Ala Val Ile Val Gly Ser Asp Pro Asp Leu Thr Lys Glu Arg Pro Leu Phe

Ile Asp Gly His Leu Arg Glu Val Gly Leu Thr Phe His Leu Leu Lys 260 265 270

Glu Met Ile Ser Ala Ala Gln Thr Ile Leu Pro Asp Ser Glu Gly Ala

245

Ala	Phe 290	Ser	Pro	Leu	Gly	Ile 295	Ser	Asp	Trp	Asn	Ser 300	Ile	Phe	Trp	Ile		
Ala 305	His	Pro	Gly	Gly	Pro 310	Ala	Ile	Leu	Asp	Gln 315	Val	Glu	Leu	Lys	Leu 320		
Gly	Leu	Lys	Glu	Glu 325	Lys	Met	Arg	Ala	Thr 330	Arg	His	Val	Leu	Ser 335	Glu		
Tyr	Gly	Asn	Met 340	Ser	Ser	Ala	Cys	Val 345	Leu	Phe	Ile	Met	Asp 350	Glu	Met		
Arg	Lys	Lys 355	Ser	Ala	Glu	Glu	Gly 360	Ala	Ala	Thr	Thr	Gly 365	Glu	Gly	Leu		
Asp	Trp 370	Gly	Val	Leu	Phe	Gly 375	Phe	Gly	Pro	Gly	Leu 380	Thr	Val	Glu	Thr		
Val 385	Val	Leu	His	Ser	Leu 390	Pro	Thr	Thr	Val	Ser 395	Val	Ala	Asn				
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gcto	tgaa	act a	agtg	gatco	cc c	eggge	ctgca	a gga	atto	gat	tgga	atga	ctc	gaaca	agctat	1	120
ggc	catga	att a	actta	agta	cc a	catg	taact	t gag	gacti	gca	atg	gacaa	agt a	actta	attatc	1	180
ctac	caaco	cta a	actto	cttt	gt to	gtgti	ttaag	g tc	caaa	aagt	tate	gcgt	gcg (	gcca	gaatcc	2	240
atca	aaaat	tgt (	gtage	catti	tg ti	tcaa	acaca	a tg	cccti	taa	ccc	gtata	agt (	gttai	tgagtt	3	300
ggta	actco	cag a	acaat	tacat	tt aa	agaga	atata	a tt	gggta	atgc	att	gttg	tgt a	atca	tataca	3	360
acad	catgo	ctc a	agtta	atgto	ca g	tatca	acaa	t cti	tcct	cttc	caa	acaca	aat	tcta	attctc	4	120

Asp Val Pro Gly Leu Ile Ser Lys Asn Ile Glu Lys Ala Leu Thr Gln 275 280 285

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<212> PRT

<213> lavendula

<400> 32

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Thr Thr Ile Thr His Gly Gly Ser His Arg Leu Pro Pro Gly Pro Arg
35 40 45

Gly Phe Pro Ile Val Gly Ala Leu Pro Leu Leu Gly Asp Met Pro His 50 55 60

Val Ala Leu Ala Lys Met Ala Lys Thr Tyr Gly Pro Ile Ile Tyr Leu 65 70 75 80

Lys Val Gly Ala Trp Gly Met Ala Val Ala Ser Thr Pro Ala Ser Ala 85 90 95

Arg Ala Phe Leu Lys Thr Leu Asp Thr Asn Phe Ser Asp Arg Pro Pro 100 105 110

Asn Ala Gly Ala Thr Ile Leu Ala Tyr Asn Ala Glu Asp Met Val Phe 115 120 125

Ala Arg Tyr Gly Pro Lys Trp Arg Leu Leu Arg Lys Leu Thr Asn Leu 130 135 140

His Met Leu Gly Asn His Ala Leu Asp Gly Trp Ala Ser Val Arg Ser 145 150 155 160

Ser Glu Leu Gly Tyr Met Leu His Ala Arg His Asp Ala Thr Arg His 165 170 175

Gly Glu Pro Val Val Leu Pro Glu Met Leu Met Tyr Ala Val Gly Asn 180 185 190

Met Leu Gly Gln Val Ile Leu Ser Arg Ile Phe Glu Lys Lys Gly 195 200 205

Lys Glu Val Asn Glu Leu Lys Asp Met Val Val Glu Leu Met Thr Ser 210 215 220

Ala Gly Tyr Phe Asn Ile Gly Asp Phe Ile Pro Trp Leu Ala Trp Met 225 230 235 240

Asp Leu Gln Gly Ile Glu Ser Gly Met Lys Lys Leu His Asn Lys Phe 245 250 255

Asp Lys Leu Ile Gly Lys Met Ile Glu Asp His Leu Lys Ser Ala His
260 265 270

Ile Arg Lys Ala Lys Pro Asp Leu Leu Asp Cys Leu Leu Ala Asn Arg 275 280 285

Asp Ser Ser Asp Ala Glu Lys Leu Thr Ser Thr Asn Val Lys Ala Leu 290 295 300

Leu Leu Asn Leu Phe Thr Ala Gly Thr Asp Thr Ser Ser Ser Ile Ile Glu Trp Ala Leu Ala Glu Met Ile Lys Asn Pro Thr Ile Leu Asn Arg Ala His Gln Glu Met Asp Arg Val Val Gly Arg Thr Arg Arg Leu Val Glu Ser Asp Ile Pro Asn Leu Pro Tyr Leu Arg Ala Ile Cys Lys Glu Thr Tyr Arg Lys His Pro Ser Thr Pro Leu Asn Leu Pro Arg Ile Ala Ser Glu Pro Cys Val Val Asp Gly Tyr Tyr Ile Pro Lys Asn Thr Arg Leu Ser Val Asn Ile Trp Ala Ile Gly Arg Asp Pro Asp Val Trp Glu Asn Pro Leu Asp Phe Asn Pro Asp Arg Phe Leu Ser Gly Lys Asn Glu Arg Ile Asp Pro Arg Gly Asn His Phe Glu Leu Ile Pro Phe Gly Ala Gly Arg Arg Ile Cys Ala Gly Ala Arg Met Gly Met Val Leu Val Glu Tyr Ile Leu Gly Thr Leu Val His Ala Phe Glu Trp Glu Leu Pro Ala Gly Ala Gly Ala Gly Thr Ala Glu Leu Asn Met Asp His Val Phe Gly Leu Ala Leu Gln Lys Ala Val Pro Leu Thr Ala Met Leu Thr Pro Arg Leu Pro Ser His Cys Tyr Ala Pro